

SEQUENCE LISTING

5

(1) GENERAL INFORMATION:

- (i) APPLICANT: GETZENBERG, Robert H.
- 10 (ii) TITLE OF INVENTION: BLADDER NUCLEAR MATRIX PROTEINS,  
POLYNUCLEOTIDE SEQUENCES ENCODING THEM, AND THEIR USE
- (iii) NUMBER OF SEQUENCES: 4
- 15 (iv) CORRESPONDENCE ADDRESS:  
(A) ADDRESSEE: Foley & Lardner  
(B) STREET: 3000 K Street, N.W., Suite 500  
(C) CITY: Washington  
(D) STATE: D.C.  
20 (E) COUNTRY: USA  
(F) ZIP: 20007-5109
- (v) COMPUTER READABLE FORM:  
(A) MEDIUM TYPE: Floppy disk  
25 (B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:  
30 (A) APPLICATION NUMBER:  
(B) FILING DATE:  
(C) CLASSIFICATION:
- 35 (vii) PRIOR APPLICATION DATA:  
(A) APPLICATION NUMBER US 08/742,850  
(B) FILING DATE: 01-NOV-1996  
(A) APPLICATION NUMBER: US 60/006,626  
(B) FILING DATE: 03-NOV-1995
- 40 (viii) ATTORNEY/AGENT INFORMATION:  
(A) NAME: BENT, Stephen A.  
(B) REGISTRATION NUMBER: 29,768  
(C) REFERENCE/DOCKET NUMBER: 76333/111
- 45 (ix) TELECOMMUNICATION INFORMATION:  
(A) TELEPHONE: (202)672-5300  
(B) TELEFAX: (202)672-5399  
(C) TELEX: 904136

50 (2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

0986657-053001

- (A) LENGTH: 6 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

5

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Leu Ala Lys Ile Val Leu  
1 5

10

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

15

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Glu Ile Ser Gln Leu Asn Ala Gly  
1 5

20

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

25

30

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Val Tyr Glu Asp Ile Met Gln Lys  
1 5

35

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 12 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

40

45

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Ser Leu Asp Leu Asp Leu Ile Ile Ala Glu Val Lys  
1 5 10

50

00666027-053001